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	<210> <211> <212> <213>	63 256 DNA Glycine max	
nds.	<400>	63	
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	acgacccaaa	a tgcttcaaag ttgatacacc atgcatgcat aacttgggga gcgtaccaa	ag 180
	tggtgaacca	a tgccataccc atgagcctcc tccaagacat tcaatgggtt ggggagaca	at 240
	cttctctctc	c ccttga	256
	<210> <211> <212> <213>	64 273 DNA Glycine max	
	<220> <221> <222>	unsure (4),(7)(9),(14)(16),(19),(24),(29),(38)(39) (48),(61),(68),(94),(127)(128),(131),(133),(250), (252),(271)	′

	<223>	unsure at all n locations	
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	ngtgaggntg	ggttaaatca aacgaaccgc tacncagcta gctaggtgca caaagccgaa	120
	cggttgnnag	ngnctgttga aatgcttgct ttagtgccaa ggtactcatt ccaagtcact	180
	gccttacaaa	ggggaggctt atttgggccc actagcttcg catgtggaca tatctccaca	240
	ttcggagggn	cnctacataa atacgcactg naa	273
tanii tanii tanii	<210> <211> <212> <213> <400>	65 263 DNA Glycine max	
und tend to	ctagtgaaag	ttctctagca aaagtcatgg gagaggtaga cccagctttc atccaagacc	60
denn med	cacaacacag	gccaaagttc tctaccatac aacctgaagc gttcctgtga tagatctctc	120
	tccaataacc	aaccacacac tttcagattc atcttccatt gaaaacttag tgcaggagat	180
j	agggagtgca	tgcaaggagt ggggtttctt ccaagtaaca aaccatgggg tgcccctcac	240
	tctaagacaa	aacattgaga tag	263
He Health Charles	<210> <211> <212> <213>	66 248 DNA Glycine max	
	<400>		60
		gcccataget tacctgatte teacgeatgg teteactete aacceaacga	120
		gtctcattca atgatgatgc atcatcatca tcattcatac ccatcataga	180
		ccaaatgcca tggaacaaat aggccatgca tgtgagaaat ggggtgcttt	240
	ccaattgaag	g aaccatggca tacccttttg tgttattgaa gatgtagaag aagaggctaa	248
	aaggctct		210
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	<222>	unsure (58)(60) unsure at all n locations	
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	ccttcactct	cagaageeta eegageeeae eeegtgeaeg tteaacaeaa geaeeetgae	120
	ttaaactccc	tacaagaact ccccgagtct tacacttgga cacaccatag ccatgatgat	180
	catactcctg	cagcttccaa cgagagtgtc cccgttattg atctcaacga cccaaatgct	240
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u u	cctgacccgg	g attataaaga cccgacccaa gaagatccgg atactatacc catcatagat	120
j.	ctctcatgct	tagaccatga cacaacaagt tggaggaagc ttgcaaggat tggggtttgt	180
	ttcgtttggt	t caaccatggg gttccattga cccttttgaa tgagcttcaa gagctggcca	240
	aagaactct	t ctctttgtcc tttgaggtga aaga	274
	<210> <211> <212> <213>	69 262 DNA Glycine max	
		a gcagcgaagt taatgtgcct tatgttggct tcccttggta ttcccaagga	60
		a geagegaage edaegegeet trij 55	
		ic ccgagttgcc cggatccgga tcgggccatg ggtctggccg cgcacacgga	
		c ctcacaatcc tgcaccaaaa caatgtcaat gggcttcagg ttctcaagga	
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	aggggaagg	gg tgggtggcgg tg	

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atcgatttga	acgacccaaa tgctccaaac ctcataggcc atgcatgcaa aacatggggt	180
	tggtgaacca tggcatcccc acgagcctct tcagtgacat tcagagggct	240
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	a agattaccac aaatactgtg atcccgtcaa gcgctatgat gaagccatga	180
aaaagctagt	t gggaaagctg atgtggctga tgttggattc tctgggtatt acaaaggaag	240
acctgaaat	g ggc	253
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	g toataattag tototaatto taattagooa tacattgaac acaccagoac	
	eg taagtggtat ttgtteeaca caggtaeact atteetteae teteagaage	
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actccccga	ag	
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		DNA Glycine max	
	<222>	unsure (152) unsure at all n locations	
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	tgcaattgaa	ctcttacccg acttgtccgg anccggatcg ggccatgggt ctggccgccc	180
	acaccgactc	caccettete acaateettt accaaaacaa cataageggg ttgcaggtte	240
	accgaaaagg	cggcgg	256
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h			
z, - E	<220> <221>	unsure	
	<222>	(128), (130), (212), (216), (238), (240), (244)(245), (248)(249)	
nyli:	<223>	unsure at all n locations	
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ı.L		gaagccatga aaaagctagt gggaaagctg atgtggctga tgttggattc	60
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		n tgcaattgaa ctcttacccg acttgtccgg atccggatcg ggccatgggt	180
	ctggccgcc	c acaccgaact ccaccctctc anaatnttta ccaaaacaaa atgggggngn	240
	tgcnngtnna	a cgg	253
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	<400>	75	60
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	gacgaggac	a aggacgacac catccagctt acaaccaaca ggctttaccc gttgcatgtg	180

tacctcacac	ctagaggaag g	aaatgagca t	cacatttat	ttggtctctg	gtctgtgagc	240
						245
atatg						
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	t gattaagcgt					
	a cacaagaaac					
	a tccaaattct					
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gcttatccc	t ggttcaggct	ttaggcgagg	g atgtgagtt	c aatctatg		200
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	actgggacgc	gcccgacgaa tggaggccag agaggtatct ggacgggagc ttcgaagtcg	180
	ctgataagta	caagaccatg gcattcggcg gaggaaggac ggactgtgcg ggaagcatcc	240
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19955	<210> <211> <212> <213>	80 263 DNA Glycine max	
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		a ttcggagccg gatcacttcc cccagtacca gcggttccag gattaccagt	240
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	tccttgaca <sup>.</sup>	t gggctcatct att	203
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		ac tgactggcga tcaaatttcc atgctaatct gggatagcat tattgagaca	
		a cattagttac tactgaatgg gctatgtatg aacttgctaa agacaaaact	
		cc gtcttcatga ggagctccaa tatgtatgtg gacatgaaaa tgttatcgtt	276
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gagaccatta						240
cttgc						245
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cggtg						245